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Substitute for form 1449A/B/PTO				Complete If Known	
				Application Number	10/692002
				Filing Date	October 24, 2003
				First Named Inventor	Mike West
				Art Unit	1644
				Examiner Name	Not Yet Assigned
Sheet	1	of	2	Attorney Docket Number	DU-P02-002

U.S. PATENT DOCUMENTS					
Examiner Initials*	Cite No. <sup>1</sup>	Document Number	Publication Date	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear
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FOREIGN PATENT DOCUMENTS					
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NON PATENT LITERATURE DOCUMENTS					
Examiner Initials	Cite No. <sup>1</sup>	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published.			
CA	SELLKE, T. et al., Calibration of pvalues for testing precise null hypotheses, <i>The American Statistician</i> , 55, 62-71, (2001)				T <sup>2</sup>
CB	BREIMAN, L., Statistical Modeling: The two cultures (with discussion), <i>Statistical Science</i> , 16, 199-225 (2001)				
CC	OSBORNE, B.G., Applications of near infrared reflectance spectroscopy to compositional analysis of biscuits and biscuit doughs, <i>J. Sci. Food Agric.</i> , 35, 99-105 (1984)				
CD	BROWN, P.J., et al., The choice of variables in multivariate regression: A non-conjugate Bayesian decision theory approach, <i>Biometrika</i> , 86, 635-648 (1999).				
CE	LI, C. et al., Model-based analysis of oligonucleotide arrays: Expression index computation and outlier detection, <i>Proc. Natl. Acad. Sci.</i> , 98, 31-36 (2001)				
CF	EARLY BREAST CANCER TRIALISTS' COLLABORATIVE GROUP, Polychemotherapy for early breast cancer: an overview of the randomized trials, <i>Lancet</i> , 352:930-942 (2001)				
CG	WEST, M. et al., Predicting the clinical status of human breast cancer by using gene expression profiles, <i>Proc. Natl. Acad. Sci. USA</i> 98, 11462-11467 (2001)				
CH	SPANG, R., et al., Prediction and uncertainty in the analysis of gene expression profiles, <i>In Silico Biol.</i> 2, 0033 (2002)				
CI	VAN T VEER, L.J., et al., Gene expression profiling predicts clinical outcome of breast cancer, <i>Nature</i> 415, 530-536 (2002)				
CJ	VAN DE VIJVER, M.J. et al., A gene-expression signature as a predictor of survival in breast cancer, <i>N. Engl. J. Med.</i> 347, 1999-2009 (2002)				
CK	HUANG, E., et al., Gene expression predictors of breast cancer outcomes., <i>Lancet</i> in press (2003)				
CL	POMEROY, S.L., et al., Prediction of central nervous system embryonal tumour outcome based on gene expression, <i>Nature</i> 415, 436-442 (2002)				
CM	ALIZADEH, A.A., et al., Distinct types of diffuse large B-cell lymphoma identified by gene expression profiling, <i>Nature</i> 403, 503-511 (2000)				
CN	ROSENWALD, A., et al., The use of molecular profiling to predict survival after chemotherapy for diffuse large-B-cell lymphoma.				

Examiner Signature	/Jerry Lin/	Date Considered	05/17/2006
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CO	BHATTACHARJEE, A., et al., Classification of human lung carcinomas by mRNA expression profiling reveals distinct adenocarcinoma subclasses, Proc. Natl. Acad. Sci. USA 98, 13790-13795 (2001)
CP	RAMASWAMY, S., et al., Multiclass cancer diagnosis using tumor gene expression signatures, Proc. Natl. Acad. Sci. 98, 15149-15154 (2001)
CQ	GOLUB, T.R., et al., Molecular classification of cancer: class discovery and class prediction by gene expression monitoring, Science 286, 531-537 (1999)
CR	SHIPP, M.A., et al., Diffuse large B-cell lymphoma outcome prediction by gene expression profiling and supervised machine learning, Nat. Med. 8, 68-74 (2002)
CS	YEOH, E.-J., et al., Classification, subtype discovery, and prediction of outcome in pediatric acute lymphoblastic leukemia by gene expression profiling, Cancer Cell 1, 133-143 (2002)
CT	CHENG, J., et al., Unique Features of Breast Cancer Res. Treat. 2000;69:213-23)
CU	MITTRA, I., et al., A Meta-analysis of reported correlations between prognostic factors in breast cancer: does axillary lymph node metastasis represent biology or chronology, Eur.J.Cancer 1991;27:1574-83
CV	MC GUIRE, W.L., Prognostic factors for recurrence and survival in human breast cancer, Breast Cancer Res Treat. 1987;10:5-9
CW	TANDON, A.K., et al., HER-2/neu oncogene protein and prognosis in breast cancer, J Clin. Oncol. 1989;7:1120-6
CX	KASS, R.E., et al., Bayes' factors, J. Am. Stat. Assoc. 80, 773-705 (1985)
CY	HOETING, J., et al., Bayesian model averaging, A tutorial, Statistical Science, 14(4), 382-417 (1999)
CZ	CLYDE, M., Bayesian Statistics 6. Bernardo J.M. (ed.), pp. 157-185 (Oxford University Press, 1999)
CA1	JATOI, I., Significance of axillary lymph node metastasis in primary breast cancer, J. Clin. Oncol. 17, 2334-2340 (1999)
CB1	PHILOSOPHOV, L., et al., Medical Diagnostic Decision Rules Based on Mutually Dependent Diagnostic Factors, Comput. Biol. Med., Vol. 27, No. 4, 329-347, 1997.
CC1	LIAO, S., et al., Appropriate medical data categorization for data mining classification techniques, Med. Inform. Vol. 27, No. 1, 59-67 (2002).
CD1	CHAPMAN, W., et al., A Comparison of Classification Algorithms to Automatically Identify Chest X-Ray Reports That Support Pneumonia, Journal of Biomedical Informatics, 34, 4-14 (2001).
CF1	DAWSON, K., et al., A Bayesian approach to the identification of panmictic populations and the assignment of individuals", Genet. Res. Camb., 78, 59-77 (2001).
CF1	Copy of International Search Report dated July 1, 2004 from corresponding application no. PCT/US03/33946, citing references PHILOSOPHOV, LIAO, CHAPMAN and DAWSON.
CG1	SORLIE, T., et al., Gene expression patterns of breast carcinomas distinguish tumor subclasses with clinical implications, PNAS, Vol. 98, No. 19, 10869-10874 (2001)
OH1	O'HAEGELEER et al. Bioinformatics, Volume 16, pages 767-726 (2000)

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